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[SMART, a simple modular architecture research tool: identification of signaling ...](#) - ► [pnas.org](#) - [Full View](#)

J Schultz, F Milpetz, P Bork, CP ... - Proceedings of the ..., 1998 - National Acad Sciences

... Recent reports (43, 44) **identify** the rostral cerebellar malformation gene product (rcm) and similar homologues ... As the DEATH domain of UNC-5 is not **annotated** in databases, this is ... in **cell** signaling (7). In accordance with this, the SMART alignment **database** contains several ...

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SG Megason, SE Fraser - Mechanisms of development, 2003 - Elsevier

... image sets and manage the large amount of data through the use of a **database**. ... all these dimensions with sufficient spatial and temporal resolution for tracking **cells** is a ... Principle among these is achieving sufficient spatial and temporal resolution for **identifying** and tracking **cells** ...

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[... 2DE **database** for spot selection, automated **annotation**, and data analysis](#) - ► [uark.edu](#) [PDF]

L Malmstrom, J Malmstrom, G Marko-Varga, G ... - Journal of Proteome ..., 2002 - pubs.acs.org

... Hence, **identifying** and filtering interesting data is necessary because of the large amount ... Our research team uses the **database** to investigate new hypothesis within human pulmonary ... Today, the 2DDB retrieves information about **annotated** proteins from Internet databases such ...

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[Transitive functional **annotation** by shortest-path analysis of gene expression data](#) - ► [nih.gov](#) - [Full View](#)

X Zhou, MCJ Kao, WH Wong - Proceedings of the National ..., 2002 - National Acad Sciences

... We check the GO process classification of the transitive genes on the SP to **identify** the L0 and L1 matches, as discussed in ... The 51 novel genes predicted with the conservative rule for which no biological process **annotation** is available in either **database** are listed in ...

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[MIPS Arabidopsisthaliana **Database** \(MaTDB\): an integrated biological knowledge ...](#) - ► [nih.gov](#)

H Schoof, P Zaccaria, H Gundlach, K Lemcke, S ... - Nucleic acids ..., 2002 - Oxford Univ Press

... Thus, manual **annotation** by experts for certain protein families should be available in a whole-genome **database**. ... structure of the phenotype catalog was designed to aid researchers observing a phenotype to query the **database** and **identify** characterized mutants with a ...

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[Molecular immunology databases and data repositories](#)

V Brusic, J Zelezniukow, N Petrovsky - Journal of immunological methods, 2000 - Elsevier

... the blind panel study were subjected to statistical analysis to **identify** possible antibody ... This is an example of a specialised **database** providing information related to a ... Users can extract fully **annotated** sequence information, pairwise comparisons, multiple sequence alignments ...

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[Database tools for integrating and searching membrane property data correlated ...](#)

JS Mirsky, PM Nadkarni, MD Healy, PL Miller, GM ... - Journal of neuroscience ..., 1996 - Elsevier

... to put only positive data in the **database** (data which affirms the presence of a property in a **cell**

compartment). ... Thus, gatekeeping is kept to a minimum; the functions of the **database** managers are to ensure that the data are formatted properly and **annotated** appropriately. ...

[Cited by 27](#) - [Related articles](#) - [All 4 versions](#)

[PDF] [Helping data suppliers and consumers negotiate the details: a **database** view ...](#)

A Rosenthal, E Sciore - submitted to Conference on Cooperative Information ..., 1999 - Citeseer

... We now try to capture the requirements implied by the above scenarios, and to **identify** the major functions of a tool that might meet them. ... We assume there is an **annotation** facility for attaching information to various granules of a **database** (eg, a table, column, **cell**, or ...

[Cited by 2](#) - [Related articles](#) - [View as HTML](#) - [All 4 versions](#)

[Models and tools for collaborative **annotation**](#) - [arxiv.org](#) [PDF]

X Ma, H Lee, S Bird, K Maeda - Arxiv preprint cs/0204004, 2002 - arxiv.org

... and **identify** particular vocalizations of interest, noting their start and end times and classifying the call types. ... The following table shows statistics for $K \times$ for the test **database**. ... Consider a boolean $n \times n$ matrix, where n is the number of anchors in an **annotation** graph. ...

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[PDBsum: a Web-based **database** of summaries and analyses of all PDB structures](#)

RA Laskowski, EG Hutchinson, AD Michie, AC ... - Trends in Biochemical ..., 1997 - Elsevier

... our own CATH classification of protein structural families and the EC, PDB Enzyme Structures

Database. ... by the SITE records in the PDB file, while the small red dots **identify** the residues ... 3).

Further details on all the **annotations** are provided by the links (not shown here) that ...

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